Attorney Doc' No.: BGI-132CP

APPENDIX A: DNA SEQUENCES

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>RXA00178-upstream

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Attorne Jocket No.: BGI-132CP

>RXA00178

>RXA00178-downstream
TAAGTAGAAGCAGCAAAAAACCA

>RXA00226-upstream

 $\tt CCGCCTGCGGTGTCAGCGGAGCGCCCCGGCGTCTGAAAACTGCACTTGGTGAGAACGTGATGTTCATGATGTCACCTGTTCCATCGGTACGGAGGGGC$

>RXA00226

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CAGCCCGCAAGCAGGGGAGCCGACGTCATCATCGCGGACCCTGCCAGACGCGCCGGGGGC
AAGCGCATTACAAATCCGGCACAGCTCCTGCCACCTCTGCCTTCCGCTTCTCGACGCCTGG
ATCAACCAACCACTCGCCGTTAAATGTGCCCCCGGCCTTGATTTTTCGGAATGGCCAGGT
CTCGTCAGTATTGCCAGCGTTGATGGAGGCGTGAAAGAAGCATGCCTCTACACTACGGAT
CTGGCAGATGGGGAAACTCGCGAAGCTATCGTGATCAAAGATGGGCTCATTGACCGCATC
ACCAACTTTGAAGACGATGCCACGGGACAAGACCTTGCGGCTGCACCTGGTGAGTTCATC
ATCGACCCAGACGGTGCCATCGTGCGCCCCGGGTTTGGTTCGCCACTATGCAGTGCGTGAG
CAGCTGTGGATGTTGGATGAGCGGATCGCATACCTTACGGCAATCGGATTCCAGAGGGT
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GCACATGATGCGGGGGCGGTTGAAATTTTAGTGCGTGGTGTTTGATGTTGATCCTGATCAG
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GGCAGCCGAGGGGTTGCATTGATTTTGTGGTCCTCGCGAGCGCCC

>RXA00226-downstream
TAAAGCCGATGCAAATAAAATTG

>RXA00231

Attorney Jocket No.: BGI-132CP

>RXA00231-downstream TAGCATCTGCCCCTTTACAAATC

>RXA00249

CAGACCCGCCGCCACCTGTCCGTATCGGCTCTGCTGGGCGTACGCACGGTGATCCTCGCA GTCAACAAATTGACCTTGTTGATTACAGCGAAGAAGTCTTCCGCAACATTGAAAAGGAA TTCGTTGGCCTGGCATCTGCACTTGATGTCACAGACACCCACGTTGTTCCAATCTCTGCG CTCAAGGGCGACAACGTTGCAGAACCTTCCACCCACATGGATTGGTACACCGGACCAACC GTGCTGGAAATCCTGGAAAACGTAGAAGTTTCCCACGGCCGTGCACACGACCTGGGCTTC CGCTTCCCAATCCAGTACGTCATCCGCGAGCACGCCACCGACTACCGTGGCTACGCCGGC ACCCAGGTCACCCACATCGATTCCGCTGACGGATCCCTCCAGACCGCATCAGTTGGAGAA GCCGTTGTCCTGCGCCTAGCCCAGGAAATCGACCTCATCCGCGGCGAACTCATCGCTGGC GAAGACCGCCCAGAATCCGTTCGCTCCTTCAACGCCACTGTTGTTGGCTTGGCCGATCGC ACCATCAAACCAGGTGCAGCAGTCAAGGTTCGCTACGGCACCGAGCTGGTCCGCGGACGC ACCTACGGCCTCAACGACATCGCACACGTGCGCATCGACGTTGCAGGCGAACTCGAAGTT GAAGATTACGCTGCCCGCGCCCCATCGGATCCTTCCTCCTCATCGACCAATCCTCCGGC GATACCCTCGCAGCTGGCTTGGTTGGCCACCGCCTACGCAATAACTGGTCGATC

>RXA00249-downstream TAGACCAGTGTCTTAGGCAAGAC

>RXA00277-upstream

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>RXA00277

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>RXA00299-upstream

Attorney Jocket No.: BGI-132CP

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>RXA00299

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CGCGCTTTACGACGCGACCCTCAGTCCGCCGAGAACTTCCCGTCCGACGTTGTCGAGCTG
AACTCTTACCTCACCGGCCGTTCCCGTCTCCCAGGGGTTAACGCAATTCCAGGCAAGGGC
ACCAACGTACCGCTGTACATCTTGGGTTCATCCCTCTTTGGTGCACAAATTGGCAGCACAG
TTGGGTATGCCTTATTCCTTCGCATCCCACTTCGCACCAACTCACCTTGAGCACGCGGTG
CAAACCTACCGGGATAACTACCAGCCTTCAGAGCAGCATCCTGAGCCTTATGTCATTGCG
GCCGTCAATGTCACCGCATCTGATTCCACTGAACAAGCCCACGATGATTTCTACAAGGTA
GCGCGTGCACGCGTGAAGAACATGGCATTGCGTGGCCGACAAGTTACTGATGAGCAACTT
GATGAACTCATGGATTCACCAGCTGCTCGCCAAATTGTCGACATGCTTCACTACACCGCT
ATAGGCACTGGATCCCGAAGTTAAAGAATACCTAGACGGTTTTGTAAAGACGGCACAGGCT
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>RXA00299-downstream TAGTACCGATGGGCCGGTAGACA

>RXA00332-upstream

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>RXA00332

>RXA00332-downstream
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>RXA00354

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TTTTATCTCGAGTCCAACAACGGTTTGTATGCAAGCCGTGGTTTCCGTGAGGCTTCTAAG
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ATGTTCTGGGGCGCGAGCCTTGATCGTGACGATGTGAACAAGATCAGTTACATCTTCAAT
TCTCAGGAAGATTTGGACGCAGCGCGTGAGGCGTTCCCTAACCTGGAGCACACCACGTGG
GGTGGTCAGACGGGTGCGTTGTTCGGCACGATCGGTGTCTCTCAACAAGAAGATCGGC
GTGGATCGCCTGCTGAAGTACCTGAACGCAGATCGCCCAAACACCATTGCGTTCGGCGAC
AGCGATGAGGATCTCTCCCTATTTGAGGCGAGCGCTTACGGCGTCGCGATGGGCGAGGCC
ACCGAATCGCTCAAGGCTGCTGCTGACCTGGTCACGGATGCTGTTGGGCAGGACGCTTG
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>RXA00354-downstream TGACCCCATCAAAGAACTTCCCA

>RXA00372-upstream

Attorne, Jocket No.: BGI-132CP

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>RXA00372

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>RXA00372-downstream TAGTTTTCCTATGCAGTTATCTC

>RXA00470-upstream

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>RXA00470

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>RXA00470-downstream TAGTTGGTCGTACGCGCGTGTCT

>RXA00471-upstream

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>RXA00471

>RXA00471-downstream TAATGGCGGCTAAAAAGAGTGGC

>RXA00499

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>RXA00500-upstream

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>RXA00500

>RXA00500-downstream TGAGTGAACAATTCGAGCTACGG

Attorney bocket No.: BGI-132CP

>RXA00501-upstream

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>RXA00501

>RXA00501-downstream TAAACCATGATCGTTTTGGGAAT

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>RXA00502

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>RXA00502-downstream
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>RXA00566-upstream

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CTTTAAAGGAGAGAATAAGATTATGAGCAGCGGATTCCAA

>RXA00566

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GACGACACCCTCTGCAAACGACATCATGGCGCAGCTCCTTGTCCTCGAAGGCATGGACCCA
GACCGCGATATCACCCTGTACATCAACTCACCTGGTGGATCCTTCACCGCGTTGATGGCA
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GCATCCGCAGCCGCAGTTCTTCTTGCAGCCGGTGCACCAGGTAAGCGCGCTGTTCTTCCT
AACTCCCGCGTGCTGATCCACCAGCCAGCAACCCAGGGCACCCAGGGTCAGGTTTCTGAC
CTCGAGATCCAGGCAGCTGAAATCGAGCGCATCGTTTGATGGAAACCACCTTGGCT
GAGCACACCGGCAAGACCGCGGAGCAGTCGTTCGATTACCGCAAGATCCTC
ACCGCTGAGGAAGCACTCGAGTATGGCATCGTTGACCAGGTCTTCGATTACCGCAAGCTC

AAGCGC

>RXA00566-downstream TAGAGTTTTTTAAAGATTCGGGT

>RXA00567-upstream

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>RXA00567

>RXA00567-downstream TAGGCGCACGGAAAACTTTAAAG

>RXA00569-upstream

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>RXA00569

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>RXA00594-upstream

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>RXA00594

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>RXA00612

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>RXA00615

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>RXA00615-downstream TGAAAATCAGGCCAGACAGTGGG

Attorne, Jocket No.: BGI-132CP

>RXA00621-upstream

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>RXA00621

>RXA00621-downstream TGAAGTACGCGTGCTTCCTTTCG

>RXA00622-upstream

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>RXA00622

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>RXA00622-downstream TAAAACTTGTTCGACTAGCGAAC

>RXA00636-upstream
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>RXA00636

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>RXA00639

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>RXA00640-upstream

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>RXA00640

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>RXA00641-upstream
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>RXA00641

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>RXA00644-upstream

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>RXA00650-upstream

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>RXA00650

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>RXA00658-upstream

>RXA00658

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>RXA00659

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>RXA00663

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>RXA00675

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>RXA00689-upstream

>RXA00689

>RXA00689-downstream
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>RXA00715-upstream

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>RXA00744-upstream

>RXA00744

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>RXA00756-upstream

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>RXA00772-upstream

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>RXA00773

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>RXA00778

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>RXA00787-upstream

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>RXA00787

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>RXA00791

>RXA00793-upstream

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>RXA00793

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>RXA00797-upstream

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>RXA00797

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>RXA00820

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>RXA00820-downstream TAAAAGCTTGCTTCTCGACGCAA

>RXA00833-upstream

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>RXA00833

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>RXA00844-upstream

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>RXA00844

>RXA00844-downstream TAGAATATGCTGATCTCCCTGCT

>RXA00857

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>RXA00866-upstream

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>RXA00866

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>RXA00888-upstream

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>RXA00888

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>RXA00903

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>RXA00905

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Attorne, locket No.: BGI-132CP

>RXA00906-upstream

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>RXA00906

>RXA00906-downstream
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>RXA00907-upstream

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>RXA00912-upstream

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>RXA00912-downstream TAAATGACCCTGCAACTATTCAC

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>RXA00961

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>RXA00964-upstream

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>RXA00977-upstream

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Attorne; locket No.: BGI-132CP

>RXA00977

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>RXA00983-upstream

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>RXA01018

Attorney locket No.: BGI-132CP

>RXA01022

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>RXA01055-upstream

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>RXA01055

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>RXA01056

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>RXA01056-downstream
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>RXA01057-upstream

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>RXA01059-upstream

>RXA01059

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>RXA01073-upstream

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>RXA01082-upstream

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>RXA01082

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>RXA01090-upstream

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>RXA01113

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>RXA01117

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>RXA01120-upstream

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>RXA01224-upstream

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>RXA01126

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>RXA01147-upstream

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>RXA01151

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>RXA01161-upstream

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>RXA01181

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>RXA01224-upstream

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>RXA01232-upstream

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>RXA01236-upstream

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>RXA01250

>RXA01254-upstream

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>RXA01254

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>RXA01257-upstream

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>RXA01257

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>RXA01270-upstream

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>RXA01288-upstream .

>RXA01288

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>RXA01288-downstream TAGTGCTTTTCGACGAGCCCTCC

>RXA01302-upstream

>RXA01302

>RXA01303-upstream

>RXA01303

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>RXA01307-upstream

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>RXA01307

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>RXA01309-downstream
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>RXA01324-upstream

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>RXA01340-upstream

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>RXA01340

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>RXA01354-upstream

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>RXA01354

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>RXA01354-downstream
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>RXA01358-upstream

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>RXA01358

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>RXA01385-upstream

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>RXA01385

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>RXA01412-upstream

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>RXA01412

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>RXA01412-downstream
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>RXA01426-upstream

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>RXA01427-upstream

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>RXA01427-downstream TAGCAGTAAACTTCTTTGAATAC

>RXA01428-upstream

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>RXA01728-upstream

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≯RXA01728-downstream TAAATAATGAGTAAAAAGTCTGT

>RXA01795-upstream

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>RXA01795

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>RXA01802-upstream

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>RXA01828-upstream

>RXA01828

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>RXA01829

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>RXA01829-downstream TGAACCGCTCGATTCGAATCACA

>RXA01838

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>RXA01848-upstream

>RXA01848

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>RXA01869-upstream

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>RXA01869

>RXA01869-downstream TAGGTTTCTGGGTGGAGAAGGAT

>RXA01885-upstream

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>RXA01885

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>RXA01914-upstream

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>RXA01914

>RXA01918-upstream

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>RXA01918

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>RXA01932

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>RXA01933-upstream

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>RXA01933

>RXA01933-downstream TGACACCTCAACCACTGATTTTG

>RXA01934-upstream

>RXA01934

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>RXA01967-upstream

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>RXA01967

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>RXA01967-downstream
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>RXA01971-upstream

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>RXA01971

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>RXA01993

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>RXA01994-upstream

>RXA01994

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>RXA01997-upstream

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>RXA01997

>RXA02000-upstream

>RXA02000

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>RXA02010-upstream

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>RXA02010

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>RXA02012

>RXA02016-upstream

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>RXA02016

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>RXA02017-upstream

>RXA02017



>RXA02017-downstream TAGGACACACTATGTCAAACTTT

>RXA02018

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>RXA02048

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>RXA02048-downstream
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Attorney locket No.: BGI-132CP

>RXA02052-upstream

>RXA02052

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GTCCGCCACGAGGCTATTTATGATCAGCTTTCAGACATCGCACCAACCGTGATGTCCGAA
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GATGAGATGGACAAACTGATCTCAGACTTGGACACCCGTGCCACAGAGCTTGGTGAAGAG
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TCCGAACTGTGGGTTGATTCACCAGCAGTTCGGCAAGGTAGAACTTACGAGTTTGAAGAC
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>RXA02064-upstream

>RXA02064

>RXA02064-downstream
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>RXA02083-upstream

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>RXA02083

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>RXA02092-upstream

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>RXA02092

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>RXA02098

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>RXA02101-upstream

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>RXA02101

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>RXA02105-upstream

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>RXA02118-upstream

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>RXA02118

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>RXA02120

>RXA02120-downstream TAAATTGCTTGTCGACGCCTAGT

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>RXA02126

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>RXA02192

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>RXA02214-upstream

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>RXA02214

>RXA02214-downstream TAACCCTGAAGGCGATTTAAGGG

>RXA02215

>RXA02215-downstream
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>RXA02264-upstream

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>RXA02264

>RXA02265-upstream

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>RXA02265

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>RXA02274

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>RXA02275

>RXA02275-downstream TGATGGATCTTGACGCTGATTTT

>RXA02276-upstream

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>RXA02276

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>RXA02277

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>RXA02278-upstream

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>RXA02278

>RXA02278-downstream TAGACGCCGTCGAGAAATCGAAG

>RXA02316-upstream

Attorne; locket No.: BGI-132CP

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>RXA02317

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>RXA02334

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Attorne; locket No.: BGI-132CP

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>RXA02746-downstream
TAAAAACTTATGAATAATCCAGC

>RXA02813

>RXA02813-downstream TAATGGATC

>RXA02820

>RXA02828-upstream AGTCAGGATTTATG

>RXA02828

>RXA02834

>RXA02834-downstream TAATCTAGCAAAATATTAATTTG

>RXA02839

>RXA02855-upstream

ACCCCGATCCTTTGTTTTCGTGGGATCACTATTAGACTCGACTCTACCGCGCTGCAGGTT TTCCTGATACGCCTGCGGACAAAACAGAAAGGTATTTCAC

>RXA02855

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GAGTATTTGGCCTCGGTTCTGAAAGATCACGGCATGGAACCACACCTGTTCCCAGGAACC GGTTTGATGGTGGATATCGGACCAGAAGGGGACTCCCGCCTGGCGTTTCGCGCTGATATC GATGCCCTTCCGCTGCTTGAATCAACCGGCTTAGAGTTCTCTTCCACAGCCACTGGCGTT GCGCATGCCTGCGGACATGACGTGCACACGGTGATCGCTTTGGCACTTGCCTGTGCACTG AACACCATCGAACTGCCCATCGGCATTCGGGTGATTTTCCAGCCGGCAGAAGAAGTCATG ACTGGTGGCGCAACGGACGTCATTGCCCACGGTGGCCTTGATGGTGTGGATGCGATTTAC GCCATCCACGTTGAACCCAAATTGAAGGTCGGTCGCGTCGGTGTACGCGCTGGCGCGATT ACTTCTGCCTCAGATGTGATCGAAATCAGAGTCAAGGGTGAAGGAGGACATAGCGCACGT CCACACCTCTCCGCTGATGTTGTTTACGCCTTGAGCAAATTGGTCGTTGATCTTCCCGGT TTGCTGTCCAGGCGCGTCGATCCACGCACCGCACCGTGCTTGTTTTCGGCACCATCAAC GCCGGCTATGCGCCCAACGCGATCCCAGATTCCGGCATCGTGTCAGGCACCTTGCGTACA GCCGACATCTCTACCTGGCGTGACATGCGTCCGCTTATCTCTGAGCTGGTGGAACAGGTG CTCGCACCCACCGGAGTCACCCATGAACTGATCTACAATCCGGGTGTTCCACCAGTGCTT AACGACGATGTCGCCACCGCTTTGTTGGCAAGCGCAGCACGCGACATGGACACACAATCT GTTGTCCAAGCGCCGCAGTCATCCGGTGGAGAAGACTTCTCGTGGTACCTTGAACACGTC CAAAGTGACCTGGTTGTGGATGAGCGAGCCATCGGAGTTGGCGTCAGGCTCTTTGGCTCC CTTGTGCAGCAGTACAGTAGCCGATCTGAAGCTTTCTTAAATTCC

>RXA02855-downstream TAATGGGGGTAGTGTGTAGGGCT

>RXA02859

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GCACGATTGCTTATCGACGCTCCCTCCCCATCACGCCGGGGTCCGGCGTGGGCCAGCTG
GGCTCCGGCAGCGCAACGATGTGGCACCGGCGACCCCAGAATTACCAGAACCCACCGAC
AACCTGACTTCAGGTGAGGGTGGGTCGCAGAACACGCTGCTTAAGCTCGTGGTGCCCATC
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>RXA02859-downstream TAGGTGTTCTTCTTCACGACCTC

>RXA02895-upstream

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TATCAGCGTAAACGTCCGAACATGAAAGGCTAGAAAAGCC

>RXA02895

>RXN00045-upstream
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>RXN00045

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>RXN00045-downstream TAAGCACCTAAAACGCTGTTCTC

>RXN00073-upstream
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>RXN00073

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>RXN00136-upstream

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>RXN00136-downstream TAAGTGTTGAGTAATCTACTAGT

>RXN00177-upstream

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>RXN00177

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>RXN00178-upstream

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AACGGCGACGATCCAGCAACTTTGAATTAAGGAGACCAAC

>RXN00178

>RXN00178-downstream TAAGTAGAAGCAGCAAAAAACCA

>RXN00249

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>RXN00249-downstream TAGACCAGTGTCTTAGGCAAGAC

>RXN00250-upstream

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>RXN00250

>RXN00250-downstream TAATCTTTCCGGTCCGCCAGTTC

>RXN00299-upstream

>RXN00299

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>RXN00299-downstream TAGTACCGATGGGCCGGTAGACA

>RXN00343-upstream

 ${\tt TTCGGTAGAATGGGTAGGTTGTCGTGCTTGAGGTGTGGTGGATAACCACCTCTACAACACCCCCAAGCTCTGTTAGAAAAAATTGAGGAAGCAGTCTAA}$

>RXN00343

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>RXN00393-upstream

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>RXN00393

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>RXN00393-downstream TAAAACGCTTTTCGACGCTCCCC

>RXN00397-upstream

>RXN00397

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>RXN00397-downstream
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>RXN00398-upstream

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>RXN00398

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GGACTTCCCGCACTTCTTCGCTTGGTAGAAAATGGCATTAAAGTCAAAGCAACCGGATTC
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>RXN00398-downstream TAGTTTTAAGACCCGAAATGTCT

>RXN00434-upstream

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>RXN00434

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>RXN00434-downstream TAGGCAAGCGATGCCAGCACCCT

>RXN00470-upstream

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>RXN00470

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>RXN00470-downstream TAGTTGGTCGTACGCGCGTGTCT

>RXN00499-upstream

 ${\tt TGCCAACAGGGGATATGCCACTGTGTACCCCACGGCGATGATCGGTAAAATCCTCGGCGCGCAGATATTGTTCTTGCTGCTCTAAGGTGATTTTTTGGGCA}$

>RXN00499

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>RXN00499-downstream
TGATTCCTCGTGTTAGTACTAGC

>RXN00513-upstream

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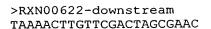
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>RXN02021-upstream

TTGGGTCGCCGAGGAGATCTAATCCTGGTTTGAGTTCAGAGTTCACAGGTTTAAGCCTAC
AAACCTTAGTTAAAACATGATGGAAGCGGTCGATTAAAAA

>RXN02021

ATGAGTGAAAACATTCGCGGAGCCCAAGCAGTTGGAATCGCAAATATCGCCATGGACGG ACCATCCTGGACACGTGGTACCCAGAACCCCAAATTTTCAACCCGGATCAGTGGGCTGAA CGCTACCCATTGGAAGTGGGCACCACACGCCTCGGAGCAAACGAACTCACCCCACGGATG CTGCAGTTGGTAAAACTGGACCAAGATCGCCTCGTCGAACAGGTAGCAGTCCGCACCGTT ATCCCCGATCTGTCTCAACCTCCAGTAGACGCGCACGATGTTTACCTGCGCCTCCACCTG CTTTCCCACCGCTGGTCCGCCCCCACGAAATGCACATGCAAAACACCTTGGAGCTGCTG TCCGACGTGGTGTGGACAAACAAGGGCCCTTGCCTTCCTGAAAACTTTGAGTGGGTGCGT GGTGCTCTGCGGTCCCGCGGACTCATCCACGTCTACTGTGTGGACCGTCTTCCCCGCATG GTCGACTATGTGGTTCCCCCTGGAGTCCGCATCTCCGAAGCAGAACGCGTGCGCCTAGGT GCATACCTTGCTCCGGGTACCTCTGTGCTGCGTGAAGGTTTCGTGTCTTTCAACTCCGGC ACCTTGGGTGCCGCAAAGGTGGAAGGCCGCCTGAGTTCCGGTGTGGTCATCGGTGAAGGT TCCGAGATTGGACTGTCTTCTACTATTCAGTCCCCGAGAGATGAACAGCGCCGCCGTTTG CCGTTGAGCATCGGCCAAAACTGCAACTTTGGTGTCAGCTCCGGAATCATCGGAGTCAGT TTCGCAGCCGATGAGGAGTTACGCACTATCGACTCCATCGAAGGCCAAGCAAATTGGTCA ATCAAGCGTGAATCCGGCTTCCATGAGCCAGTTGCCCGCCTCAAAGCT

>RXN02021-downstream TGACCCATTTTCATAACCAGTGC

RXC00110 - 5'-Region

RXC00110 - coding Region

GTGAGCAACAAGACGGCCTTTTTACTGACGGTAACAGCACGTTTGCACCTAAGGTGGATTCAATTCCC
CTCAGCGATGTGGATACCAGCGTTAGCGGTGAAGCCTCCATCGGCACGCTGATCTCCAACGCAACCTCC
CAAATGTCCAGCCTTTTCCGCGCAGAAGTTGAGCTGGCGAAGACTGAACTCGCAGGCGAAGCCAAGAAA
GCTGCCATCGGCGGCGGCGCATTCAGCGTTGCTGGCGTAATCGCACTGTACAGCTCCTTCTTCTTTTTC
TTCTTCGTCGCAGCACTGCTGAGCGAGTGGATTAAGCCTTGGGCAGCATTCCTCATCGTGTTCCTCTTC
ATGCTGGTCATCGCCGCAGCTCTCGCACTGTTCGGCTGGCGCAAGGTGAAGAAGATGGGCGCTCCGAAG
AACACCATCCAATCGGTCAACCAACTGAAGAACCTGGTCCCAGGTCAGGCATCCGAGAAGCTGGAGAAG
GCCAACAAGCGTGGCCTCTACACCTCCGCGTCCTTCCACAGCCCCGGCGCCATCACTGGCGACCAC

RXC00110 - 3'-Region
TAAAAAAGGAGACTTCGATGGCC

RXC01971 - 5'-Region

 ${\tt AGGTCTTGTTTATTTCGGCTACTGATTCAGTAGCTGCGCTCCGATAGGATTCTTAGTTTCAGTTCAGTAGCTTTGAGCCACGGCTAGAATGTGAATCCT}$

RXC01971 - coding Region

RXC01971 - 3'-Region
TAACGTGAGGTAGCTCACAGTCA

RXS00004 - 5'-Region

TTGCACTGTCATGACTGTATCCCGCGAAGAAGTGTCCCTGCCGAGCCGAACTCTGAACAATGCCTTCCGGAAGTATTTTCCAATTCCCGATGTAGGGTCA

RXS00004 - coding Region

GTGCTGACTCAATTGATTGAATCATCGATTTTCGACAACGTTGCGAGCAGGGAGTCCTCTGAATTTCTC
GGCCATGCTGCCATCGATCTACTTGCTGGCCTTGTCTATGAAAAAGCCACTCCCTATGCTCCAGATGAA
GCACTTAGAGTGGCAGTTTATGGCTATATTCGGGAGAACCTTGGATCCTCACAACTTACGGTCGCAGCT
GTAGCCGGGGCGCATAGAATCGCGGTTCGTACGTTGCATCGATTATTTGAAGGCGAAGCATACGGAGTA
GCGGAATTAATCCGACACCTCCGATTAGAGGCAGTATATGAAGACCTTCGGGATCCTCGCCTCCAGAAC
CTGACCATTTTGGCTATCGGCATGCGCCACGGCATTTCCAGCCAAGCTCATTTAACAAGACTGTTTCGC
GCTAAATATGGGGTACCGCCGGCAGAGTTTCGCCGAGGGTATATTAATAGCGCTGCT

RXS00004 - 3'-Region TGAGGGCACCGCAAGCGTGGCGC

RXS00156 - 5'-Region

 ${\tt ACCGAGAGCGTGGTACGCCTCATTTAGTTTCCTCCTATGAATCTTGATGTGGTTCATGCGTTTTTATGCAATCTAACCAAAAGTTGGTACGATCCTCAT}$

RXS00156 - coding Region

ATGAATGAACGCACATCGGATGCATTTGACGCCCTCCTTGTGCTCCTTCGGTGGTCCCGAAGGGCAC GAGGAGGTTCGTCCGTTTTTGGAGAATGTCACTCACGGAAGGGGGATTCCGCCGGAACGTCTAGATGAA $\tt GTGGCGGTTCATTACCACCACTTCGGTGGTATCAGCCCCATCAATGCGCTGAACAGGGAAATTATCGCC$ AATGTGGAAAAAGAATTGGCGTCTCGCGATCACAAGCTGCCTGTTTATTTTGGTAACCGCAACTGGAAG CCGTTTGATAATGAGGCCGCTGAACAAATGGCTGATGACGGCGTGAAAAAACGCGCTGGTGTTGGCAACT TCCGCTTGGGGTGGCTACTCCGGTTGTCGGCAGTACCAGGAAGATATTCAGGGCATGATCAAGCACCTG GAGTCTCAGGGGCAGTCGATCACGTTCACCAAGCTGCGTCAGTTCTACGATCACCCTCGTTTTGTCTCC ACCATGGCTCAATTGGTTCAGGATTCCTACGCGAAGCTTCCCGATGAGCTGCGAGATGAGGCGCGTCTG GTCTTCACCGCGCACTCCATTCCACTGACTGCGGACAATGCTGCGGGAACCCCTGAGGATGGCTCCTTG TATTCCACACAGGTCAAGGAAGCGTCAGCACTGATTGCTGAGGCTGTTGGTGTCAGATTTTGATGTG GTGTGGCAGTCCCGCTCGGGTAGCCCGCACACTCCGTGGCTGGAGCCTGACATCGTGGATCACGCAGTG GAGCTCAACGAGAAGGGTCAAAAAGCGCTCGTTGTCTGCCCTGTAGGCTTTATTTCTGATCATATGGAA GTCATTTGGGATCTTGATTCCGAGCTGATGGAAGAAGCCGAGAAGCGCAACATGGTGGTCGAGCGTGTC GCTACCGTTGGCCCCACCGATGAATTCGCAGCCCTTGTGGTTGATCTCATCGAGGAGGCAGAGCTCAAG CGCGTTATCGAGCGCCTTGGAAAGCTGCCAGCACGCGGAAGTTCCGTCAACGGCGCACCGTGTGGCGAC **GCCAAC**

RXS00156 - 3'-Region TAGGAGTGATAGTCCCTCGCAAA

RXS00166 - 5'-Region

GGCGTTTAGCGATCTTCAACATCGAGCAACCAGCGCCAGCGCTTTTACCCAAGGCAGCACGACTTATCACGATGTCCGACCTGGATATCCGGCTGAGGCC

RXS00166 - coding Region

RXS00166 - 3'-Region TAGGAGCCCTCGCCATGTACCTG

RXS00197 - 5'-Region

RXS00197 - coding Region

 $\tt GTGGCAGCCTATCTTCTTGGTGTCGTATTATTTTTCCTCGGCATCGCAGTAACCATCGCGCTTCACGAG$ TGGGGGCACTTCATCACAGCGCGCATTTTCGGAATGAAAGTGCGGCGTTTCTTCATCGGTTTCGGCCCG ACGGTGTTTGCCAAAAGACGCGGCGAAACCGTGTACGGCCTTAAAGCGATTCCGGTCGGCGGTTTTTGT GACATCGCGGGGATGACTGCCCAAGATGAACTTGATCCGGAAGACCTGCCGCGCGCCATGTATCTAAAG $\verb|CCCTGGTGGCAGCGCATAATTGTGCTTTCCGGCGGCGTGATCATGAATCTGATCGTCGGCTTTTTGGTG|\\$ CTTTACGGCGTGGCGTGAGCTCCGGAATCCCGAATCCGGATGTGGATACCACCGCGACAGTCGACACC GTTCAGTGCGTGCCGGAAACCCAAATTTCCGCAACTGAACTGTCCTCCTGCGTAGGTTCAGGCCCAGCG GGCGACGCCGGCATTGAGCACGGCGATAAGATTTTGGCCGTCAACGGCCAAGAGATGGCAAGCTTCACC CTTTTCGACGTCGACCTCCAGGTTGCCTCTGTCACCCGTCTCGCCTCTGACGGTTCAGAAATTACCGTC GGCGCGGTGGGCATGTCGAGCCTTCCACCGACCGATGTGTACAAAAAATACGGCCCAATCGAGGGTGTG GGAGCAACTGCACGTTTCACCGGCGACATGATCAGCGCCACGTGGGATGGCCTCAAAGCCTTCCCGGCG AAAATCCCAGGGGTCGTCGCATCCATCTTCGGTGCAGAACGAGATGTAGAAAGCCCCATGAGTGTGGTG GGCGCCGTACGCATCGGCGGCGAATTTGTCGAACGTTCCATGTGGGACATGTTCATGATGATGCTGGCC AGCCTGAACTTCTTCCTCGCGCTGTTTAACCTCGTGCCGCTGCCACCACTTGATGGCGGACACATTGCC GATTACACCAAACTAATGCCCGTCACCGTAGCTGTCGCAGCCTTGCTGATGACAGTGGGAGGCCTGGTC ATTGTCGCCGATGTGGTCAATCCCATCCGACTCTTTGGC

RXS00197 - 3'-Region TAACGATACGGAATTGAACTGCC

RXS00288 - 5'-Region

GGCGTGCTAAAAAAGCĀCGTCAAATACAGAATCGGCAGATTACATCGCTGAGCAGAAAAACACGGGCA TGAAACGTACCCAAGGGCTAACATCGGGGGC

RXS00288 - coding Region

RXS00288 - 3'-Region TAAAGAGGACTTGAAAAATGCAC

RXS00624 - 5'-Region

RXS00624 - coding Region

ATGTCCGGCCGTCTTCTTGTTTCAGTTTCTAGTATTTTCGACCAGACCCGATCGGCGGCTGACAGGCTCATTTCAGACCTGCGAGCCGACGGCATCGAGGTCTCATTACTTGTCGCACCCCGCATCGATGGGGACTGG



CGTCTCGCCAAAGACAAAGGGACCCTCGCGTGGATGGAACAACAACGCGAACGCGGCCACGAACTCATC
CTCAACGGTTTCGACCAAGCAGTTCAGGGACGTCGCTCAGAATTCGCCAACCTTGAACGGCACGAAGCA
CGTCTTCGCCTTACCGGTGCCATTAGGCAAATGCAGAAAATTGGCTTCGAATTCCAAATCTTTGCCCCA
CCTCGTTGGAGAATGTCAGAAGGCACCTTCGCGGTACTCCCAGAATTTGATTTCAACGTCGCCGCCTCG
ACCAGGGGATTACATAACCTCGACACCGGCGAATTCTTGGCGTGTAGAAACCTCTCCGTGGGTGAAGGT
TTTGGTGCTGCAAAATGGTGGCGCAAGAATGTCATCAAGGCTGTCACTCGTGGAGCGGAAAAAGGAAAT
ACAGTGCGCTTGTCCGCATCGGCGCGAAATCTCACCAACCCTAAAGTCGCAGCTGACTTCCGGGAAGCT
GCATTAGCTGCCTTGGATTTGGGTGCTCAGGTGCAAACCTATTCTCAGGCGGCCGCACAACTGGCC

RXS00624 - 3'-Region TAGTTGGGGAGGTTCGGGGCACC

RXS00949 - 5'-Region

GCTCAAGGATCCTTCCTGGGCAAACCAAGCAGCCCTCGCACTAGGTGCGGAACCCAGGTATGTTCACCA ATACGACTACGTACTTTAAAGGAGAGTTGAC

RXS00949 - coding Region

ATGAAGGTTTTCATCATCGGCGCTGCGGGTGGCATCGGCAATCGACTTTCCAGCCTGCTTCACGCCAGG
GGAGATGCAGTTAGCGGCATGCACCGCAATCTTGAGCAGGCCTCAAAAATCACAGACACTGGGGCAACT
GCCGTACTCGGGGATCTCATCCACAACAGCACGGAGGAGCTTGCGGAGCTTTTCCGCGGTCACGATGCC
ATCGTATTTTCTGCAGGCCCCACGGAACAGGGCAAGAAAATACCACGCTTATCGACGGCGCCGGCCTC
CGTAAAGCCGCCGACGCTGCCAGCGCGCCCAACGTTTCACGCTTCATCTTGGTCTCTGCGTTTCCGGAA
TCCTCCCGCGGGGAGAACACCACCGAGAACTTTGAGCACTATATGAAGGTGAAGAAGTCCGCCGATGTC
TACCTCAGTCACACTGACCTAGACTGGGTTATTTGTCCGACCAGGCGTGCTTCAAGATGAGGCAGGGGAT
GGTTTAGTCACTGCTGGCTTAGCGATTAATTACGGCAATGTTGCTCGCGATAATGTCGCAGCGTTCATT
GATGAAGCTCTGCATCAACCGCAGTTGTCAAAGATCATTGTTGAACTCACCGACGGTTCAACTCCGGTG
GCGGAAGCCGTAGAACGCCTCATCAAG

RXS00949 - 3'-Region TAAAGACGAAAAGAGGGAGAATG

RXS01000 - 5'-Region

 $\tt CTTTCTATGCCTACGCGGATGTTTCCGTGATCATTCTGGAAATCCTCATCGTGGTGATTGTCATTGAAGTATCTCCAACGCACTTCGAAAGAGGCTGGT$

RXS01000 - coding Region

RXS01000 - 3'-Region TGACGCTCCACCAAGCATCCGCA

RXS01002 - 5'-Region

GACTGCTGATACCGCACAGGATGAAATCACTCGTTACGGCGAGATCCTGAAGAAGTTCTCCAACTAATT TCCCTGTTTCCAATACTCAAGGTGTGCGCAT

RXS01002 - coding Region

ATGAATTCTGATGCTTCGGCTACCACCAACTCCTGGGCTATCAACTTCGACCATGTGTCGGTGACGTATCCCAATGGGACGAAAGCCCTCGATGATGTTTCCCTCACCATCAATCCCGGTGAGATGGTTGCCATCGTGGGTCTGTCAGGATCGGGTAAATCCACGCTGATTCGCACGATCAACGGTCTTGTCCGCGCTACGGAAGGC

ACCGTGACGGTGGGGCCGCATCAGATCAACACCTTGAAGGGGAAAGCACTGCGTGATGCCCGTGGGCAG
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CGCTTCGCGCACACAGCGTGGTGGCGTAACCTCCTCGGGTTTCCCACGGAGCACAAGCAGATTGCT
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CAGAAACAGCGCGTTGCTATTGCGCGCGCCTTATCGCAAGATCCGTCTGTCATGCTGGCAGATGAGCCT
GTGGCAAGCCTTGATCCGCCAACCGCGCATTCCGTGATGCGCGATCTAGAAAACATCAACAACGTGGAA
GGCCTCACCGTGTTGGTGAACTTGCACTTGATTGATTTGGCTCGTCAATACACCACAAGGCTTGTGGGT
TTGCGTGCCGGCAAGCTGGTCTATGACGGTCCTATCTCTGAGGCCACCGATAAAGACTTTGAAGCTATC
TATGGTCGCCCCCATCCAGGCTAAAGACCTGCTAGGTGATCGCGCA

RXS01002 - 3'-Region TGACCACGCCTTCTTCTACACTT

RXS01003 - 5'-Region

RXS01003 - coding Region

RXS01003 - 3'-Region TGAGCACCTTAACCTCTCACCGC

RXS01114 - 5'-Region

TTCGGTGGAAGATATCCGCAAGCTGGTGCTGCCGCAGCTTTTAGAAACGGCTCAAGCAATTTCGACAGA TCTCTCTGCACTCTAAATTAAGGATCAAAAA

RXS01114 - coding Region

ATGAACCCTCAAGATATTGTCATCTGTTCCCCATTGCGCACCCCAGTTGGTGCTTACGGCGGATCCTTC ACCGGCGTCCCTGTTGAAGAATTGGCCACCGTGATCAACGCGATCGTTGAGGCAACCGGCATCACC GGCGACGATGTGGACGATCTGATCCTCGGCCAGGCATCCCCCAACGGTGCGGCTCCAGCACTGGGCCGT GTTGTTGCTCTAGATTCCAAGCTTGGCCAAAACGTTCCAGGCATGCAGCTTGATCGCCGCTGTGGTTCC GGCCTGCAGGCAATCGTCACCGCTGCTGCACACGTTGCATCCGGCGCTGCTGATCTGATCATCGCAGGT GGCGCAGAATCCATGAGCCGCGTTGAGTACACCGTGTCCGGCGATATCCGTTGGGGTGTCAAGGGCGGC GACATGCAGCTTCGTGACCGCCTTGCAGAAGCACGCGAAACCGCTGGCGGACGCAACCACCCGATCCCT GGTGGCATGATCGAGACCGCTGAGAACCTGCGTCGCGAATACGGCATCTCCCGCGAGGAGCAGGACAAG ATCTCCGCAGCGTCCCAGCAGCGTTGGGGCCAAGGCTGCTGATGCGGGGCTTTTCGACGACGAGATCGTG CCAGTCACCGTCCCTGCCAAGAAGCGCGGCCAGGAGCCAACCATCGTTTCTCGAGACGAGCATGGTCGA CCAGGAACAACCGTCGAAAAGCTTGCTGCTTTGCGCCCCATCATGGGCCGCCAGGATGCGGAAGCAACC GTCACCGCTGGCAACGCGTCCGGCCAAAATGATGGCGCTGCTGCCGTCATCGTGACCACTCGCGCCAAG ACCATGGGTATTGGACCTGTTCCTGCCACCAAGAAGGTCCTGGATCGTTTGGGCCTTACCCTGGAGGAC TCTTGGGAAGATGAGCGCGTCAACCCACTGGGTTCCGGTATTTCCATGGGACACCCAGTCGGTGCCACC GGTGCTCGCATGGCAGTAACCTTGGCTCACCGCATGCAGCGTGAAAACACTCAGTACGGACTGGCCACC ATGTGCATCGGTGGCGGCCAGGGTCTTGCAGCTGTCTTTGAAAAGGAGAAC

RXS01114 - 3'-Region TAAAAATGGCTATTTTGCACAGC



GTATCGGCCTATCCGCCGGCCATCATCGCAGCGGCTCTCGTAGGAATTTGCGCGGGAGTTTTGCCCCAT
AATTTTGAACCCTCGCGAATATTTATGGGCGATTCCGGCTCCATGCTCATCGGCCTGCTGTTGGCTGCA
GCATCGACCTCAGCGTCAGGAAAAATCAACATGAGCCTGTATGGCGCAGCTGATTTTATCGCATTGATC
TCACCCATCATCGTTGTTCTCGCCGCCGTGGCCATCCCACTGCTCGACCTCGTGATGGCAGTGGTTAGG
CGCGTGGGCAGGGGAGCATCACCCTTTTCCCCGGACAAAATGCATCTGCACCACCGACTGCTGTCCATC
GGACACACCCATAGGCGCGTGGTCCTAGTGCTCTACACCTGGGCGAGCGCCGTGGCATTCGGCGCAGTG
AGCTTCTCCGTCGTTCCGCCACTGTTTGCCACCGGATCGAGCATCTGTGGCATCCTCATCGCCGTCGCT
GTCACAGCCGTGCCAGTGATGAAAAGCCGGCGAGCCGCCAAACTTGAT

RXS01205 - 3'-Region TAAGTGATTGTCACTTTGGATTG

RXS01223 - 5'-Region

GCGGGAGTGGACGGCCCAGCCATCGGGGCGACAATCACGGGAGTTTTCAACGTGTCAAGGATGCTC ATGGCACCATCCTAGGCGCGCCTGCCATAGG

RXS01223 - coding Region

RXS01223 - 3'-Region TAGATTGCTAGAGATTCCCGCAC

RXS01269 - 5'-Region .

 ${\tt GATTATCTCTGCGCCGATTCAGCTGGCAATAGCAGCAGTTGTATTGAGAGCTCATGGTCGTCCGATTCTATTTCGTCAACCACGACCTGGGAAAGACGGT}$

RXS01269 - coding Region

RXS01269 - 3'-Region TAGCAGGTAAAAATTTTACTTTC

RXS01421 - 5'-Region

TTGATGCACGTGCAGAAATCGTCGGCGGTCCGTGGCACCCATCTGTTAAGGGAGACTCGGTTACTGCAGGGATCCTGCGAGATCGAGTAAACGCCTAAAG

RXS01421 - coding Region

RXS01421 - 3'-Region TAAAGCTGTGGCTGCGGAATGCG

RXS01491 - 5'-Region

 ${\tt AACGGTGTGAAACACATTGAAGTGTCGTTGGGTTATCCGCGCGAATGGGAGCATTGGTCTGGGCAGCATGTGGGCCATATCCAGTGATGGAGGTGGACA}\\$

RXS01491 - coding Region

RXS01491 - 3'-Region TGAACTGGATGGAGAGGATACCT

RXS01572 - 5'-Region

RXS01572 - coding Region

RXS01572 - 3'-Region TAGGTTTACTGAAGTTCAGACTT

RXS01642 - 5'-Region

 ${\tt TTGGAATCACTGTGTCTTGCAGCTAGTCTGCCACTCGCACTGCATTTGCTGGATGTATTTGGAATGCTGCGGGGTCTTGATATTGGATTCGGTGGATAAG}$

RXS01642 - coding Region

ATGCGCAGACTCATCGCGGTTAGCTTGGCCGCTCTGTTTATGTTGGCTTCCACTCCAGCGACGAGGGCA CACCTTTCGCAGTCATTATCCCAAGCTCACCAACTAGCAACTGGCGCCGGTGTGATGGTGGCAGTCATC GACACCGGAGTATCCCTGCATCCACGTCTGCCCCACTTAATTCCCGGCGGTGATTTCGTGGGCGCCCAC CAAAGCCCCGATGTGCCAGGTGAACTTATCGATTGCGACGGCCACGGCACCATCGTCGCCGGAATCATC GCCTCCCAAGGAAACCCCGGCACCGGCTGGCCATATGACGGCAGCTCCGATCCTTATATCGGTGTCGCC CCAGATTCCGGAATCATCTCCATTAAACAAACCAGCTCATATGTGCGTACTCGTGAAGATTCCAACGTC GGAACGCTGAGCACCCTGGCGGAATCCATCCACCGAGCTCTCGATTCCGGTGCCCACGTGATCAATATT TCCGTGGTGTCCTGTTTGCCCCAATCACCCGACGAGGCCGCATCGTTCCAGCCTCTGACGGATGCTCTT AACAGAGCAGAACTTCAAGGGGTGATAGTGGTGGCAGCAGCAGGAAACCTCGGGCAGGATTGTCCAGTT GGATCTACCGTTTATCCTGCACATTCAGACACTGTGCTCTCTGTGTCGGCACGTTTTGATTCTCACACG CTTGCAGAATATTCCATGCCTGGCAACCAACAAATCCTCTCTGCACCAAGCCACATTCAGGCTGGTCTA TCACCGCGTGGCGACGGCTTCGCCAGCCACATGATCACCACCGCTGGCGAAAGCCCCTTCGAGGGCACC AGTTTTGCCGCTCCAGTTGTCAGCGCCACAGCTGCACTGCTTCGCCAGCATTTTCCCTTTGCCACACCC TATGAAATTCGTGCACGAATCTTCAACAGCATCGACCCTGCAAGAGGCGCTATTGATCCCTACCTGGCA CTTACTCAAGAAATCTATCCCACCACTCCCCTGGTTCATGAGATCGCACTAAGTGTTCCCACGCCGCCG GATGATTCTCCACGGGAGCGGGCATCCTAGTTACCGCAATCATTGTTGGGTTGCTCGCAGTGTTAGCT

RXS01642 - 3'-Region



RXS01902 - 5'-Region

GGCTGGTCCGAAGTGCAAACCTTCAACACCGGCACCTACGGTGACAACTGGAACTTCCTCTTCTTCGGCGACACCCAGCTGTACAACACCCACTCCAACC

RXS01902 - coding Region

GTGCAGAAGAGTCCAGAACTGGGCAAACAACCTGGAACGGCGCCCACCAATCGAAAACCCAGGAACC TCCTTCATCCTCTCCGCGGGTGATCAGGCAAACCACTCCAGCTGGGACGAGCACTCCGCATACATCTCC CCAGAAACCCTGCGCAACTACCGTCTGGCCGTGAACAATGGAAACCACGACCAGTACAACTACGACGCC TACAACGCGATGTACCCACGCCCTAACCAGGTCGATGAGAACTACTTCTTCGAGTACAACAATGCACTC TTCCTGTCCCTGGACTCCAACGACTACTTGGACATCGACGACGACATCGCATTCCTTCGCGACACCGTC GCAGCACACGGTGACGACAAGGACTGGATCGTCCTGACCTACCACCATTCCACTTTCTCCCAGGCCTAC CACATGATGACGCTCGCATTAAGTACCAGCGCGAACGCCTCACCCCAGTGATCTCTGAACTGAACGTT GACTTGGTTCTCGGTGGACACGACCACATCTACACCCGCTCCCACCTGATGAACGGCTTCACCCCAGTC GATGCAGGCCGCGAAGCAGTTGTCGGTGAAACTCTGAACCCTAAGGCCGGCGAAGTTGTTTACCTTGCA ACCAACTCTTCCTCAGGCTCCAAGTTCTACGACTTCTACGACTTCCAGCTCGGCCAGCGTTACGACACC GGACTGGATTTCCAGGAAACCGTCGATCAGAAGAAGATCCGCACCTACACCGCAGTCTGGAACCAGGAC CAGGTTCAGGACTACACCAACGTTGAACTGACCCCAGAAGGCCTGACTGTGACCACTAAGGACGCAGTC TCCGGCGAGCTGGTTGACCAGTTCACCCTGAGCAAGCAGGACCGCGACGAAGAATCTGAAGTCCCAGTT GAAGATGACAAGGACGGACAACGCGACCGGCTCCTCCAACCTTGGTCTAGCTGCTATCTTGGCTCCA GTTCTGGCCATCTTCGGTTCGTCGGTGGACTCTTTGTTGGCGGCGGCTCCCTCGCTGAGTTCTTTGCC AACCTCGGCGTGAAGATGCCTTTC

RXS01902 - 3'-Region TAATACTGTCTGAGATTCAAGCA

RXS02453 - 5'-Region

AACCAACAAAGGTCATCTCAACCGGCTTAAGAAAATTCTGCCAGCTTTCTGCTGATTGAATCGTGCCAGCTCAGGGCATATCTCACCTAAAGTAAACACC

RXS02453 - coding Region

RXS02453 - 3'-Region

TAAATGAATTGATTATTTTAGGC

RXS02474 - 5'-Region

RXS02474 - coding Region

ATGAGCAAAGTTGCAATGGTTACCGGTGGTGCACAAGGCATCGGTCGTGGAATTTCAGAGAAGCTGGCA
GCAGATGGTTTCGATATTGCCGTAGCCGACCTGCCACAACAGGAAGAACAAGCTGCAGAGACCATCAAG
TTGATTGAAGCTGCAGGTCAAAAGGCTGTATTCGTTGGATTAGATGTCACCGATAAGGCTAATTTCGAC
AGTGCAATTGATGAGGCAGCAGAGAAACTTGGCGGCTTCGATGTGCTAGTAAACAACGCCGGCATCGCA
CAAATTAAGCCACTTCTGGAAGTCACCGAAGAAGACCTAAAGCAGATCTACTCCGTGAACGTTTTTAGC
GTATTTTTTGGTATTCAAGCAGCATCCCGAAAGTTCGATGAGCTTGGCGTAAAAGGCAAGATCATCAAC
GCTGCATCAATCGCTGCTATCCAAGGTTTCCCAATCTTGAGCGCCTACTCCACCACCAAATTCGCGGTT
CGTGGCCTCACCCAGGCTGCTGCGCAAGAACTCGCACCCAAGGGTCACACCGTGAATGCCTACGCACCT
GGCATCGTGGGCACCGGAATGTGGGAGCAAATCGATGCCGAGCTTTCCAAGATCAACGGCAAGCCAATC
GGTGAGAACTTCAAGGAGTACTCCTCCTCAATCGCATTTGGGCCGACCATCAGTACCTGAGGATGTAGCC

GGTCTGGTTTCGTTCCTGGCTTCTGAAAACTCCAACTACATCACCGGACAGGTCATGCTTGTCGACGGCGGCATGCTCTACAAC

RXS02474 - 3'-Region TAGGGGTTGCTTTCCCGCACTCA

RXS02485 - 5'-Region

RXS02485 - coding Region

TTGGATTCATCGCTAGCCCAGGAAATCGCCGCGATCGACGCGTCGAACTCGATTCGGAAGTCACTTTC GCCGATCTGACGACCCTCCGCATCGGCGGAAAACCCCGCAGCGCGTACGTTGCCAGACCACGGAGGCG CTCGTCGTGGCCGACGGCGATCTGGATGTTATTGCCGTCATCATCGAAACCGACGACGTCTCCATCAAC CTCACCGACGGTCTCCTCACCGCCGATGCAGGCGCTGTTTGGGACGATGTTGTCCACCTTTCGGTGGAT GCCGGCCTCGGTGGAATTGAATGCCTCTCCGGAATCCCCGGCTCCGCCGCGCCCCCCAGTCCAAAAC GTGGGCGCCTACGGCACGGAAGTTTCCGATGTACTCACCCGCGTCCAGCTTCTCGACCGCACCACCAC CAAGTCTCCTGGGTCGACGCCTCCGAACTCGACCTCTCTTACCGATACTCCAATCTCAAATTCACCAAC CGCGCAGTCGTCTTGGCGATCGAACTCCAGCTCCTCACCGACGGATTGTCCGCGCCGCTACGTTTTGGT GACGCCGTCCTAGAACTCCGCCGCGCCAAAGGCATGGTCGTGGAACACACCGACCACGACACCTGGTCC GCCGGATCCTTCTCACCAACCCAATCGTCGACCCAGCCCTTGCCGACGCAGTCTTTGAAAAAGTCGGC GAACCCACCATGCCCGCTTCCCAGCCGGCGATGGCAAAGAAAACTCTCCGCAGCCTGGCTCATCGAA CGCGCCGGCTTCAAAAAGGGACACCCCGGCGCAGGCGCAAAAGCCTCCCTGAGCACCAAACACACCCTC GCACTCACCAACCGTGGCGACGCCCGCGCCTCCGACCTCGTCGCATTAGCCAAAGAAATCCGCGACGGA GTCCTCGAAACCTTCGGCGTCACCCTCGTCCCAGAACCCGTCTGGATTGGAATCAGCATCGATGAC

RXS02485 - 3'-Region TGAATTTTCCGACGTCCCTGGCA

RXS02539 - 5'-Region

 ${\tt GGCTGCTAAGCGTGCGAATGTGCGCGTTGTCACAATCGTTGACCAAGTGTCACCTGACGCACAGGTAGTGCTCAGGTGGAGGTGGCCCAAAGGAGACCCA}$

RXS02539 - coding Region

ATGACTGTCTACGCAAATCCAGGAACCGAAGGCTCGATCGTTAACTATGAAAAGCGCTACGAGAACTAC ATTGGTGGCAAGTGGGTTCCACCGGTAGAGGGCCAGTACCTTGAGAACATTTCACCTGTCACTGGTGAA GTTTTCTGTGAGGTCGCACGTGGCACCGCAGCGGACGTTGCACTGGATGCTGCACATGCAGCC GCTGATGCGTGGGGCAAGACTTCTGTCGCTGAACGTGCTCTGATCCTGCACCGCATTGCGGACCGCATG GAAGAGCACCTGGAAGAAATCGCAGTTGCAGAAACCTGGGAGAACGGCAAGGCAGTCCGTGAGACTCTT GCTGCAGATATCCCACTGGCAATCGACCACTTCCGCTACTTTGCTGGCGCGATCCGTGCTCAGGAAGAT CGTTCCTCACAGATCGACCACAACACTGTTGCTTACCACTTCAACGAGCCAATCGGTGTTGTTGGTCAG ATCATTCCTTGGAACTTCCCAATCCTCATGGCTACCTGGAAGCTCGCACCGGCACTTGCTGCAGGTAAC GCGATCGTCATGAAGCCAGCTGAGCAGACCCCAGCATCCATTTTGTATCTGATTAACATCATCGGCGAT CTCATCCCAGAGGGCGTCCTCAACATCGTCAACGGACTCGGCGGTGAAGCAGGCGCTGCACTGTCCGGC TCTAATCGGATTGCCAAGATTGCTTTCACCGGTTCCACCGAGGTCGCAAGCTGATCAACCGCGCTGCA CTGTCACAGGATGACGCCTTCGCAGAGAAGGCAGTTGAAGGCTTCGCGATGTTCGCCCTCAATCAGGGT GAAGTTTGTACCTGTCCTTCCCGTGCACTTGTTCATGAGTCCATCGCTGATGAATTCCTCGAGCTTGGC GTGAAGCGAGTTCAGAACATCAAGCTGGGTAACCCACTTGATACTGAAACCATGATGGGTGCTCAGGCG TCCCAGGAGCAGATGGACAAGATCTCCTCCTACCTGAAGATCGGCCCAGAAGAAGGCGCTCAAACCCTC ACTGGTGGCAAGGTCAACAAGGTTGATGGCATGGAGAACGGTTACTACATTGAGCCAACCGTTTTCCGC GGCACCAACGACATGAGGATCTTCCGCGAGGAAATCTTCGGACCAGTCCTTTCTGTTGCTACCTTCAGC GACTTCGATGAGGCCATCCGTATTGCAAACGACACCAACTACGGCCTCGGCGCTGGTGTCTGGAGCCGT GACCAAAACACCATTTATCGTGCAGGTCGCGCAATCCAGGCTGGTCGAGTTTGGGTCAACCAGTACCAC AACTACCCAGCGCACTCCGCTTTCGGTGGATACAAGGAGTCCGGCATCGGCCGTGAGAACCACCTCATG ATGCTGAACCACTACCAGCAGACCAAGAACCTGTTGGTCTCCTACGATCCAAACCCAACCGGACTGTTC

RXS02539 - 3'-Region TGATCTAAGCGTTAAGTCCTAGA RXS02554 - 5'-Region

 ${\tt GCTTTTGAAGTGTCGCGTGTGCGGACTGAAATAGTTTCCGCTTCAACTTGGTTGCTAAGGATAGGCTCCATAAAAATAACCAAAGGCGGAAAATTTCA}$

RXS02554 - coding Region

ATGTCACACATAAGCCATCCATTGCCATCCTCGGTGCTGGCCGAGTGGGTTCTTCACTTGCCAGGTCA
GCGGTCGCCGCAGGCTATGAGGTAAAGGTTGCTGGTTCAGGTGGTCGACAAAAATCGCTCTTACCGCT
GAGATCCTTATGCCCGGCGCGGTTCCAAGCACTGCTGACCAGGCTGTAAAGGATGCAGATATTGTGTTC
TTGGCTGTTCCCCTGCATAAATTCCGCAGTGTCAATCCAGCCACTTTAGAGGGCCAAGATCGTTATTGAC
ACGATGAACCACTGGGTTCCGGTCAATGGTGAGTTGGAGGAAATTGATCAGGATCCGCGCAGCACTTCG
GAGATTATTGCGGAGTTTTTCGCGGGATCAACCATGGTGAAGTCTTTTAACCACATTGGTTATCACGAG
ATTGAGCAGGATGCGGGTACCGGGCGTGCGATTGCCTATGCCACGGATGATGTGGATGCAGGTGCCCAG
GTTGCACAGCTAATTAAGAGTTTTTGGGTTTGTTCCTTTAAATATTTGGCGCATTGGAAAACCGCCGTATT
CTGGAACCTGGCCAAGAAGCTTTCGGCGCGCCCCTTAATAAAGATTCGCGCCTAGAACTTGTTAATCAG
CGG

RXS02554 - 3'-Region TAGTACCTCGATCTTCAGCCAAC

RXS02560 - 5'-Region

 ${\tt TTGGGGCAAGCCAGCTAACGCATTTCTTGTGGAAACCGCAGACATTGAGGCCGCCCACGCGGAACTTCTAAGAGCAGTGGAATGAAATAATCCGGTGCTG}\\$

RXS02560 - coding Region

RXS02560 - 3'-Region
TAAAAGCATGATTATGGACGCCT

RXS02578 - 5'-Region

GGCAAAAATGAGGAACAGCACGCCCGCAATAATGAGGACCGTTGCAGATCGCTTCATAAAAAACAGCCCACACCTTTCCGCTAAACTCGCATGTTGAAATA

RXS02578 - coding Region

GTGACGGTCTTGTTGACGATCCCAGTTGCCGCCATCAGCTACACCTTCATCGAAGAGCCCATCAGCGGT GGACCCGGCGCGCATTCAGGCTGGGGGTCGTTAGGATTCACCATTTTTCTGGGGGTAGGTCTGGAAAA

RXS02578 - 3'-Region TGATGAATTGGCACCACGTCAAG

RXS02741 - 5'-Region

 ${\tt ACTGGTCACCTGGTTTGGTCTGCACTCTGACTCCCCTCAAAAGGGCACAATTTGGTCAATTTCCCAACCTTGTCTTTCAGTCATGGTTAGTGTGGGAACC}$

RXS02741 - coding Region

RXS02741 - 3'-Region TAAACACGCATAAAAAGATCCTG

RXS03058 - 5'-Region

ACGAGCTTCCGCTCTGCACAAGCCGCTAGAAGCCCCGCATAGCCCTAATGTAGAGCTCATGCCCATTTGGAATCACAACACCGCATATCGGCCATGGCTG

RXS03058 - coding Region

GTGTCAAAGCTCAAAGGCTCAAGATCGCTTCTCGACGTCGGCTCCGGCGATCACTCCTTCGCCGACCTG
GCCGGCCGCCAGGTCGCCATGTCGATGTCGTGGATCCTCTTATTAATACAACCTTTGAAGAATTCCAG
CCGACCCAAAGCTACGATGCCATCACGTTCATCGCGTCCCTCCATCACATGAACGCGGAAGAAGGACTT
AACAAAGCAGTCCGAATCCTCAATCCTGGCGGCAAGCTCCTCATCGTAGGCCTCGCCAAAAACAAAACC
GCCTCCGACTGGATCATCTCCGGACTACAAGCTTTTCTCTCCCGACCAATCAGCCTCATCAATAGGGAA
CAACAAATCTACCCCTTCCCTACCAAAGAACCCTCAGAGAGTCTCCACGAAATACGACAACTCACCAAG
CAGCTCCTCCCTCACCGCCGTATTCGCCGTGGAATCCACTTCCGATACCTCCTCGAGTGGACAAAGCCT

RXS03058 - 3'-Region TAAACAGCCCTATAAACCAAAAA

RXS03061 - 5'-Region

 $\tt CTGCCACCACTGGTCATTGCAGAGGACACTCTCCGTGATGGTCTTCAGGTGTTAGTCGCAGCCCTAGAGCGCGAAACCGCGCACCAGAAGGTGGGCTAAA$

RXS03061 - coding Region

GTGTCTTTGACCTTCCCAGTAATCAACCCCAGCGATGGCTCCACCATCACCGAGCTAGAAAACCACGAT TCCACCCAGTGGATGTCCGCGCTCTCTGATGCAGTTGCAGCTGGTCCTTCATGGGCTGCGAAAACTCCC CGCGAAAGATCCGTGGTACTCACCGCAATCTTCGAAGCACTGACCGAACGCGCCCAAGAACTTGCAGAG ATCATCCACCTGGAAGCTGGAAAATCCGATGCAGAAGCTCTTGGTGAAGTCGCTTATGGTGCAGAATAC TTCCGTTGGTTGCGGAAGAAGCAGTGCGCCTGCCCGGCCGCTACGGACAGTCA

RXS03119 - 5'-Region

TGGGAGGTGTCGCACCAAGTACTTTTGCGAAGCGCCATCTGACGGATTTTCAAAAGATGTATATGCTCGGTGCGGAAACCTACGAAAGGATTTTTTACCC

RXS03119 - coding Region

ATGGCTGTATACGAACTCCCAGAACTCGACTACGCATACGACGCTCTCGAGCCACACATCGTCGCTGAA
ATCATGGAGCTTGACCAGTCCAAGGACCACGCAACCTACGTTGCGGGCGCAAATGCAGCACTC

RXS03119 - 3'-Region
TAGGCACTAGAGAAGGCACGCGA

RXS03120 - 5'-Region

CTGGGCAGTTCTTGGGTACGACCACATATCCGGTCGCCTGGTTATCGAGCAGCTCACCGACCAGGAGGGCAACATCTCCTTCGACATCACCCCAGTTCTG

RXS03120 - coding Region

RXS03120 - 3'-Region TAAGCATTTTTAGTCCGTGCAAT

RXS03150 - 5'-Region

TTTAACAGAGTGCGTTTCAATGCCTGTAGTGTTCCGGCAATTTTGAATGTCGTTACGGTTACCCAAGGCTGAATTCCTGAGCTCACCTTGTACAAGATCA

RXS03150 - coding Region

RXS03150 - 3'-Region TAATCCAAACTGCACCTATATAT

RXS03218 - 5'-Region

 ${\tt TTGGTACGGGGGTTAACAACCTAGCTTTGTATCTGAAGGCTTCGGAGATTTCCTGTATACATCAACAAT}\\ {\tt TGCCCTTTAACCAGGAGTATTCTTAGCTTCT}$

RXS03218 - coding Region

RXS03218 - 3'-Region TGAGTTCTAGCAAATGGCGGCTTGTTAGTG